		<b>1</b> 9	14249	5 <b>T</b>
1531	Slb1———	9 ,	• , .	• ,
	GAACACCATTGAATGGGATTATTGGWATGACYCAGTTGT			1590
${ t GluIleA}$	${f xrgThrProLeuAsnGlyIleIle}$ GlyMetThrGlnLeuS ${f e}$	erLeuAs	spThrGlu	530
	H1			
TTGACRC	AGTACCAACGAGAGATGTTGTCGATTGTGCATAACTTGG	CAAATTO	CCTTGTTG	1650
LeuThrG	:InTyrGlnArgGluMetLeuSerIleValHisAsnLeuA	laAsnSe	erLeuLeu	<b>550</b>
ACCATTA	TAGACGATATATTGGATATTTCTAAGATTGAGGCGAATA(	GAATGA	CGGTGGAA	1710
	leAspAspIleLeuAspIleSerLysIleGluAlaAsnA			570
CAGATTG	ATTTTTCATTAAGAGGGACAGTGTTTGGTGCATTGAAAA	CGTTAG	CCGTCAAA	1770
	spPheSerLeuArgGlyThrValPheGlyAlaLeuLysTh			590
011111011	.bpi medel 2 daming dig i mi (ali medel medel e e e e e e			
CCTTTC	AAAAAAACCTAGACTTGACCTATCAATGTGATTCATCGT	TTCCAG	ΑΤΑ ΑΤΟΤΤ	1830
	SluLysAsnLeuAspLeuThrTyrGlnCysAspSerSerPl			610
ATATIEG	tunysksineukspheutiirtyrdiiicyskspberberr	ici ion.	Spasified	010
3 0000030	ATAGTTTTAGATTACGACAAGTTATTCTTAACTTGGCTG	CT & & TC (	מתא תייים א א כי	1890
TIEGIA	AspSerPheArgLeuArgGlnValIleLeu <u>AsnLeuAlaG</u>	IVASHA.	rarrerAz	630
	N Table of the control of the contro		mama a a m	4050
TTTACTA	AAAGAGGGGAAAGTTAGTGTTAGTGTGAAAAAGTCTGATA	AAATGG	IGTTAGAT	1950
PheThrL	ysGluGlyLysValSerValSerValLysLysSerAspL	ysmetva	alleuAsp	650
	TTGTTGTTAGAGGTTTGTGTTAGCGACACGGGAATAGGTA			2010
SerLysL	LeuLeuLeuGluValCysValSer <u>AspThrGly</u> IleGlyI	leGluL	ysAspLys	6 <b>70</b>
	G1			
	${f T}{f G}{f A}{f T}{f T}{f T}{f C}{f A}{f G}{f C}{f T}{f C}{f T}{f C}{f T}{f C}{f T}{f A}{f C}{f T}{f C}{f T}{f A}{f C}{f C}{f C}{f A}{f C}{f $			2070
LeuGlyL	${ t LeuIlePheAspThrPheCysGlnAlaAspGlySerThrT}$	hrArgL	ysPhe <u>Gly</u>	6 <b>90</b>
	-SIh2			
GGTACAG	GTTTAGGGTTGTCAATTTCCAAACAGTTGATACATTTAA'	TGGGTG	GAGAGATA	2130
GlvThrG	SlyLeuGlyLeuSerIleSerLysGlnLeuIleHisLeuM	etGlyG	lyGluIle	710
	G2	_	_	•
	ACTTCGGAGTATGGATCCGGRTCAAACTTTTATTTTACGG	TGTGCG	TGTCGCCA	2190
TrpValT	ThrSerGluTyrGlySerGlySerAsnPheTyrPheThrV	alCvsV	alSerPro	730
p				
 ТСТААТА	ATTAGATATACTCGACAAACCGAACAATTGTTACCATTTA	GTTCCC.	ATTATGTG	2250
	[leArgTyrThrArgGlnThrGluGlnLeuLeuProPheS			750
# CINSIII	reargiyrimiargoinimioradianeancarior med	0100111		
ው ጥ ል መመጥር 	GTATCGACTGAGCATACTCAAGAAGAACTTGATGTGTTGA	GAGATG	GAATTATA	2310
	/alSerThrGluHisThrGlnGluGluLeuAspValLeuA			770
Leurnev	alber interdats internet det die das praine da	rgaspo	TATTETTE	770
CAACMMC	GATTGATACCTATAATAGTGAGAAATATTGAAGATGCAA	C 2 MMC 2	CTGAGCCG	2370
				790
Granea	${ t GlyLeuIleProIleIleValArgAsnIleGluAspAlaT}$	пгреиг	nigiurio	7 30
CMC 3 3 3 6	na maa maa maa mmaa mmaa a mmaa a maa a a a a mmaa a a a a	A CHIMC A	CCDDCDDX	2430
	PATGATATAATTATGATTGATTCGATAGAGATTGCCAAAA			810
varrysi	t FyrAspIleIleMetIleAspSerIleGluIleAlaLysL	Asreny	rgrearea	010
		a.a.a	mamma	0400
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SerGlu	ValLysTyrIleProLeuValLeuValHisHisSerIleP	roGinL	euAsnmet	830
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ArgVal	${ t CysIleAspLeuGlyIleSerSerTyrAlaAsnThrProC}$	ysSerI	leThrAsp	850
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. •	GAAGCGATTAAGAGGAATAAATATGATGTGGTGTTGATGGATG	2784 928

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120 40	GACCCTGAGCTTTATAGTCAGCACTGTCATAGCCTTAGGGAAACACTTCTTGATCATASPProGluLeuTyrSerGlnHisCysHisSerLeuArgGluThrLeuLeuAspHis	A <i>I</i> As
180	AACCATCAAGCTACACTTATCGACACTTATGAACATGAACTAGAAAAATCCAAAAAT	TT
60	AsnHisGlnAlaThrLeuIleAspThrTyrGluHisGluLeu <mark>Glu</mark> LysSerLysAsn	Ph
240 80	AACAAAGCGTCCCAACAAGCACTTAGTGAAATAGGTACAGTTGTTATATCTGTTGCC AsnLysAlaSerGlnGlnAlaLeuSerGluIleGlyThrValValIleSerValAla	GC A1
300	GGAGACTTGTCGAAAAAAGTTGAGATTCACACAGTAGAAAATGACCCTGAGATTTTA	AI
100	GlyAspLeuSerLysLysValGluIleHisThrValGluAsnAspProGluIleLeu	Me
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540 180	CAAGTGCGAGAAATTGCTGATGTCACACGTGCTGTTGCCAAGGGGGACTTGTCACGT GINValArgGluIleAlaAspValThrArgAlaValAlaLysGlyAspLeuSerArg	AA As
600	ATTAATGTACACGCCCAGGGTGAAATCCTTCAACTTCAACGTACAATAAACACCATG	AA
200	:leAsnValHisAlaGlnGlyGluIleLeuGlnLeuGlnArgThrIleAsnThrMet	Ly
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720	GTATATTAGGAGGACAAGCGTTGATTGAAAATGTTGAAGGTATTTGGGAAGAGTTG	CT
240	GlyIleLeuGlyGlyGlnAlaLeuIleGluAsnValGluGlyIleTrpGluGluLeu	Le
780	ATAATGTCAATGCCATGGCTCTTAATTTGACTACACAAGTGAGAAATATTGCCAAT	AC
260	AspAsnValAsnAlaMetAlaLeuAsnLeuThrThrGlnValArgAsnIleAlaAsn	Th
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280	ThrThrAlaValAlaLysGlyAspLeuSerLysLysValThrAlaAspCysLysGly	Va
900	TYCTTGATTTGAAACTTACTATTAATCAAATGGTGGACCGATTACAGAATTTTGCT	GA
300	leLeuAspLeuLysLeuThrIleAsnGlnMetValAspArgLeuGlnAsnPheAla	G1
960 320	CGGTGACGACATTGTCGAGAGAGGTTGGTACTTTGGGTATTTTGGGTGGACAAGCT laValThrThrLeuSerArgGluValGlyThrLeuGlyIleLeuGlyGlyGlnAla	CT Le
1020 340	TACAGGATGTTGAAGGTGCT <u>TGG</u> AAACAGGTTACAGAAAATGTCAACCTAATGGCT danga d	AA As
1080 360	ATTTAACTAACCAAGTGAGATCTATTGCTACAGTTACTACTGCAGTTGCGCATGGT ssnLeuThrAsnGlnValArgSerIleAlaThrValThrThrAlaValAlaHisGly	AC Th
1140 380	TGTCGCAAAAGATTGATGGTCATCCCAAAGGAGAGATTTTACAATTGAAAAATACA 1euSerGlnLysIleAspGlyHisProLysGlyGluIleLeuGlnLeuLysAsnThr	GA As

ATCAACAAGATGGTGGACTC GCAGTTGTTTGCATCAGAAGTG GAAAGTGGCACAA IleAsnLysMetValAspSerLeuGlnLeuPheAlaSerGluValSerLysValAlaGln	1200 400
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·CAGTTGATACATTTAATGGGTGGAGAGATATGGGTTACTTCGGAGTATGGATCCGGRTCA GlnLeuIleHisLeuMetGlyGlyGluIleTrpValThrSerGluTyrGlySerGlySer	2160 720
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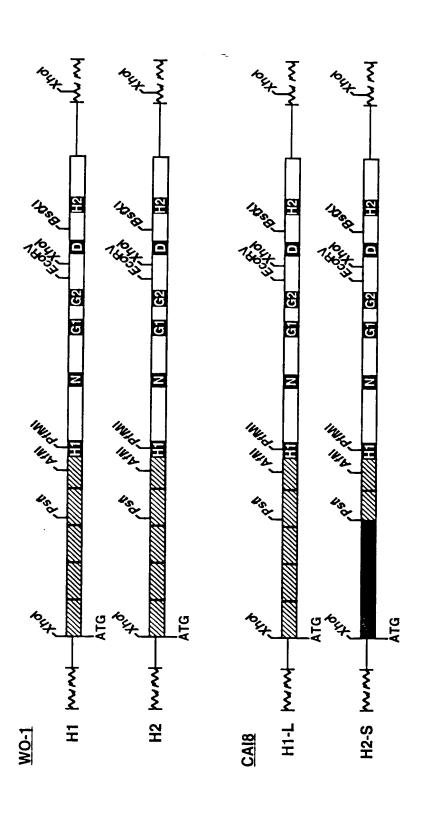


Fig. 3

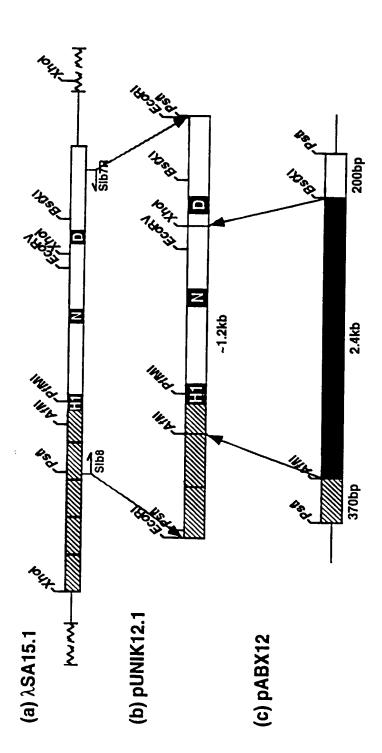


Fig. 4

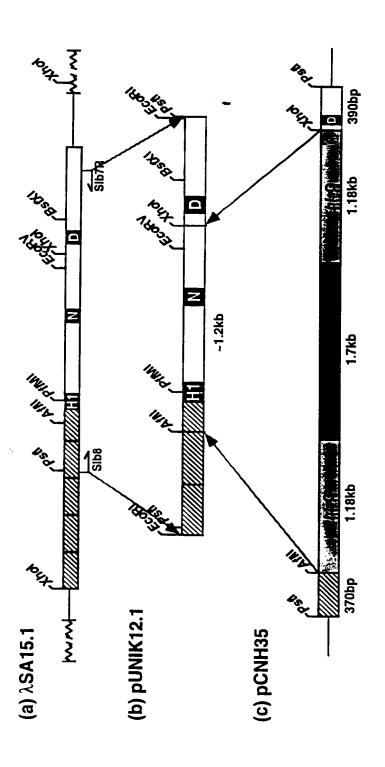


Fig. 5